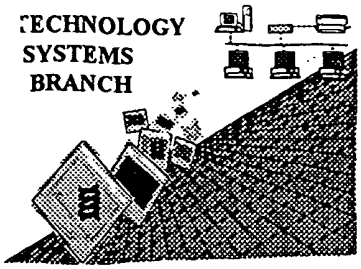
OIPE

TECHNOLOGY SYSTEMS BRANCH



0590
0124

# CRF Problem Report

**The Scientific and Technical Information Center (STIC) experienced a problem when processing the following computer readable form (CRF):**

Application Serial Number: 09/915,181
Filing Date: 7/24/2001
Date Processed by STIC: 1/26/2002

STIC Contact: **Mark Spencer, 703-308-4212**

**Nature of Problem:**

The CRF (was):

J (circle one) Damaged or Unreadable (for Unreadable, see attached)
____ Blank (no files on CRF) (see attached)
____ Empty file (filename present, but no bytes in file) (see attached)
____ Virus-infected. Virus name: _____ The STIC will not process the CRF.
____ Not saved in ASCII text
____ Sequence Listing was embedded in the file. According to Sequence Rules, submitted file should **only** be the Sequence Listing.
____ Did not contain a Sequence Listing. (see attached sample)
____ Other:

_____

**PLEASE USE THE CHECKER VERSION 3.1 PROGRAM TO REDUCE ERRORS. SEE BELOW FOR ADDRESS:**
**http:/www.uspto.gov/web/offices/pac/checker**

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.
Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.
Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<**http://www.uspto.gov/ebc/efs/downloads/documents.htm**>, **EFS Submission User Manual** - ePAVE)
2. **U.S. Postal Service**: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
3. **Hand Carry directly to:**
   U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
   Or
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Revised 01/29/2002

0570
0405

#9

OIPE

RAW SEQUENCE LISTING DATE: 04/08/2002
PATENT APPLICATION: US/09/915,181A TIME: 15:33:00

Input Set : A:\305-932610US.txt
Output Set: N:\CRF3\04082002\I915181A.raw

p.6

ENTERED

<110> APPLICANT: EDWARDS, ROBERT
BELLOCCHIO, ELIZABETH
FREMEAU, ROBERT
REIMER, RICHARD
<120> TITLE OF INVENTION: NOVEL GLUTAMATE TRANSPORTERS
<130> FILE REFERENCE: 305T-932610US
<140> CURRENT APPLICATION NUMBER: US 09/915,181A
C--> <141> CURRENT FILING DATE: 2002-03-26
<150> PRIOR APPLICATION NUMBER: US 60/220,556
<151> PRIOR FILING DATE: 2000-07-25
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<221> NAME/KEY: misc_feature
<223> OTHER INFORMATION: n is a, g, c, or t
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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/915,181A

DATE: 04/08/2002
TIME: 15:33:00

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Output Set: N:\CRF3\04082002\I915181A.raw

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 83 gccctggtgc actacagtgg agtcatcttc tacggggtct ttgcttctgg ggaaaaacag  1560
 85 gactgggctg atccagagaa tctctctgag gagaaatgtg gaatcattga ccaagatgaa  1620
 87 ttagccgagg aaacagaact caaccacgag gctttcgtaa gtcccagaaa gaagatgtct  1680
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 93 tcttaacgtg catcttcccc tcagcttaca accagaagtc tccacaccca ttgcttttcc  1860
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111 tgtcctctca gagccaaagc tgagactggc agatgaccag gagttttagg aaggaaggaa  2400
113 ggaaggaagg aaggaaggaa ggaaggaagg aaggaaggaa ggaagggttc agttgagtgt  2460
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128 <221> NAME/KEY: misc_feature
129 <223> OTHER INFORMATION: n is a, c, g, or t
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137 gtttcttttg agctaccctg gttgctcctc cctctacggt aactcgactc actccttcct   180
139 tccggacacg tctgtaggtc tcgggctcgg ggacacacgc tgacgtcgac gacaccgtag   240
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163 ctctgttcat atcctcttcc gcggttgaac cggtcagact cgtttaagtt gtgtggtacc   960
165 tcttccaaaa agtgtaggaa cggacagata cggtaataac accgtttgaa aacatcttcg  1020
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169 cgttattcat tccacccaga gaacagtcga cagggtgtgt accactactg ttagcaccat  1140
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RAW SEQUENCE LISTING DATE: 04/08/2002
PATENT APPLICATION: US/09/915,181A TIME: 15:33:00

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Output Set: N:\CRF3\04082002\I915181A.raw

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179 taggagtacc cctagagttt accgcaccct tgggagagac cttaccaaac aggggagtaa   1440
181 caaccacgtt actgtttcgt gttctgggcc cttcttaccg tcttacacaa ggagtatcgt   1500
183 cgggaccacg tgatgtcacc tcagtagaag atgccccaga aacgaagacc ccttttgtc   1560
185 ctgacccgac taggtctctt agagagactc ctctttacac cttagtaact ggttctactt   1620
187 aatcggctcc tttgtcttga gttggtgctc cgaaagcatt cagggtcttt cttctacaga   1680
189 atacctcggt ggtgggtctt aacactccag gtcttctgcc tagcctctgt tgtctctctt   1740
191 aggcggaagc tccccctcct cggtaatagg atggtcttac ttctcctgaa aagtctttgt   1800
193 agaattgcac gtagaagggg agtcgaatgt tggtcttcag aggtgtgggt aacgaaaagg   1860
195 gtatggaacc ggaaggtccc ccggtttagt gtcctttccc cctctgattt agttgttgtc   1920
197 tcttcttttt acggaagaat gtttctaccc gcatacctag aaccagagtc aattaatcta   1980
199 tcaactagta taaaaaaaac ccccccccgtt aacccgtaac cgacaactcg gaagagagtt   2040
201 ttcttgttaa ataagtcctt ctttaccgat cttcttattc ctcaccgaac aacgagttta   2100
203 tttgtgactt ctttagggag aaaccagacc tcttctcatg taccaccaac ggtggggtag   2160
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207 ccgggaacca acacggtcca cgaaatactt gtaagaataa attgagggtg tgggattata   2280
209 tcaataacat gggtaaaatg ttgattcttg taatttactg atccaaccgg gtgggttcca   2340
211 acaggagagt ctcggtttcg actctgaccg tctactggtc ctcaaaatcc ttccttcctt   2400
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215 tcccagtaaa agttactgtt tttgtttttg accttagtca accaaacacc cattaaggta   2520
217 caaaccagtt cccacacacg tacgtttgca catacacgca cacacacaca aacacacaaa   2580
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223 <211> LENGTH: 850
224 <212> TYPE: PRT
225 <213> ORGANISM: Homo sapiens
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233             20                  25                  30
235 Val Gly Asp Ser Leu Gly Ile Leu Gln Arg Lys Leu Asp Gly Thr Asn
236         35                  40                  45
238 Glu Glu Gly Asp Ala Ile Glu Leu Ser Glu Glu Gly Arg Pro Val Gln
239     50                  55                  60
241 Thr Ser Arg Ala Arg Ala Pro Val Cys Asp Cys Ser Cys Cys Gly Ile
242 65                  70                  75                  80
244 Pro Lys Arg Tyr Ile Ile Ala Val Met Ser Gly Leu Gly Phe Cys Ile
245                 85                  90                  95
247 Ser Phe Gly Ile Arg Cys Asn Leu Gly Val Ala Ile Val Glu Met Val
248             100                 105                 110
250 Asn Asn Ser Thr Val Tyr Val Asp Gly Lys Pro Glu Ile Gln Thr Ala
251         115                 120                 125
253 Gln Phe Asn Trp Asp Pro Glu Thr Val Gly Arg Ala Asn Ser Leu Ile
254     130                 135                 140
256 His Gly Ser Phe Phe Trp Gly Tyr Ile Val Thr Gln Ile Pro Gly Gly
257 145                 150                 155                 160
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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/915,181A

DATE: 04/08/2002
TIME: 15:33:00

Input Set : A:\305-932610US.txt
Output Set: N:\CRF3\04082002\I915181A.raw

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262 Phe Leu Thr Ser Thr Leu Asn Met Phe Ile Pro Ser Ala Ala Arg Val
263             180                 185                 190
265 His Tyr Gly Cys Val Met Cys Val Arg Ile Leu Gln Gly Leu Val Glu
266         195                 200                 205
268 Gly Val Thr Tyr Pro Ala Cys His Gly Met Trp Ser Lys Trp Ala Pro
269     210                 215                 220
271 Pro Leu Glu Arg Ser Arg Leu Ala Thr Thr Ser Phe Cys Gly Ser Tyr
272 225                 230                 235                 240
274 Ala Gly Ala Val Val Ala Met Pro Leu Ala Gly Val Leu Val Gln Tyr
275                 245                 250                 255
277 Ile Gly Trp Ala Ser Ala Phe Tyr Ile Tyr Gly Met Phe Gly Ile Ile
278             260                 265                 270
280 Trp Tyr Met Phe Trp Leu Leu Leu Gln Ala Tyr Glu Cys Pro Ala Val
281         275                 280                 285
283 His Pro Thr Ile Ser Asn Glu Glu Arg Thr Tyr Ile Glu Thr Ser Ile
284     290                 295                 300
286 Gly Glu Gly Ala Asn Leu Ala Ser Leu Ser Lys Phe Asn Thr Pro Trp
287 305                 310                 315                 320
289 Arg Arg Phe Phe Thr Ser Leu Pro Val Tyr Ala Ile Ile Val Ala Asn
290                 325                 330                 335
292 Phe Cys Arg Ser Trp Thr Phe Tyr Leu Leu Leu Ile Ser Gln Pro Ala
293             340                 345                 350
295 Tyr Phe Glu Glu Val Phe Gly Phe Ala Ile Ser Lys Val Gly Leu Leu
296         355                 360                 365
298 Ser Ala Val Pro His Met Val Met Thr Ile Val Val Pro Ile Gly Gly
299     370                 375                 380
301 Gln Leu Ala Asp Tyr Leu Arg Ser Arg Lys Ile Leu Thr Thr Thr Ala
302 385                 390                 395                 400
304 Val Arg Lys Ile Met Asn Cys Gly Gly Phe Gly Met Glu Ala Thr Leu
305                 405                 410                 415
307 Leu Leu Val Val Gly Phe Ser His Thr Lys Gly Val Ala Ile Ser Phe
308             420                 425                 430
310 Leu Val Leu Ala Val Gly Phe Ser Gly Phe Ala Ile Ser Gly Phe Asn
311         435                 440                 445
313 Val Asn His Leu Asp Ile Ala Pro Arg Tyr Ala Ser Ile Leu Met Gly
314     450                 455                 460
316 Ile Ser Asn Gly Val Gly Thr Leu Ser Gly Met Val Cys Pro Leu Ile
317 465                 470                 475                 480
319 Val Gly Ala Met Thr Lys His Lys Thr Arg Glu Glu Trp Gln Asn Val
320                 485                 490                 495
322 Phe Leu Ile Ala Ala Leu Val His Tyr Ser Gly Val Ile Phe Tyr Gly
323             500                 505                 510
325 Val Phe Ala Ser Gly Glu Lys Gln Asp Trp Ala Asp Pro Glu Asn Leu
326         515                 520                 525
328 Ser Glu Glu Lys Cys Gly Ile Ile Asp Gln Asp Glu Leu Ala Glu Glu
329     530                 535                 540
331 Thr Glu Leu Asn His Glu Ala Phe Val Ser Pro Arg Lys Lys Met Ser
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RAW SEQUENCE LISTING DATE: 04/08/2002
PATENT APPLICATION: US/09/915,181A TIME: 15:33:00

Input Set : A:\305-932610US.txt
Output Set: N:\CRF3\04082002\I915181A.raw

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335                 565                 570                 575
337 Gln Gln Arg Glu Ser Ala Phe Glu Gly Glu Glu Pro Leu Ser Tyr Gln
338             580                 585                 590
340 Asn Glu Glu Asp Phe Ser Glu Thr Ser Arg Ala Ser Ser Pro Gln Leu
341         595                 600                 605
343 Thr Thr Arg Ser Leu His Thr His Cys Phe Ser His Thr Leu Ala Phe
344     610                 615                 620
346 Gln Gly Ala Lys Ser Gln Glu Arg Gly Arg Leu Asn Gln Gln Gln Arg
347 625                 630                 635                 640
349 Arg Lys Met Pro Ser Tyr Lys Asp Gly Arg Met Asp Leu Gly Leu Ser
350                 645                 650                 655
352 Leu Asp Ser Ser Tyr Phe Phe Trp Gly Gly Gln Leu Gly Ile Gly Cys
353             660                 665                 670
355 Ala Phe Ser Gln Lys Asn Asn Leu Phe Arg Lys Lys Trp Leu Glu Glu
356         675                 680                 685
358 Gly Val Ala Cys Cys Ser Asn Lys His Arg Asn Pro Ser Leu Val Trp
359     690                 695                 700
361 Arg Arg Val His Gly Gly Cys His Pro Ile Ser Lys Asp Ile His Val
362 705                 710                 715                 720
364 Glu Asp Asn Leu Cys Asn Leu Met Lys Gly Ile Thr His Gly Gly Pro
365                 725                 730                 735
367 Trp Leu Cys Gln Val Leu Tyr Glu His Ser Tyr Leu Thr Pro Thr Pro
368             740                 745                 750
370 Tyr Ser Tyr Cys Thr His Phe Thr Thr Lys Asn Ile Lys Leu Gly Trp
371         755                 760                 765
373 Pro Thr Gln Gly Cys Pro Leu Arg Ala Lys Ala Glu Thr Gly Arg Pro
374     770                 775                 780
376 Gly Val Leu Gly Arg Lys Glu Gly Arg Lys Glu Gly Arg Lys Glu Gly
377 785                 790                 795                 800
379 Arg Lys Glu Gly Arg Lys Gly Ser Val Glu Cys Arg Val Ile Phe Asn
380                 805                 810                 815
382 Asp Lys Asn Lys Asn Trp Asn Gln Leu Val Cys Gly Phe His Val Trp
383             820                 825                 830
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388 Val Phe
389     850
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392 <211> LENGTH: 582
393 <212> TYPE: PRT
394 <213> ORGANISM: Rattus rattus
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402             20                  25                  30
404 Lys Lys Gln Asp Asn Arg Glu Thr Ile Glu Leu Thr Glu Asp Gly Lys
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RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/09/915,181A

DATE: 04/08/2002
TIME: 15:33:01

Input Set : A:\305-932610US.txt
Output Set: N:\CRF3\04082002\I915181A.raw

Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 2582,2587,2590,2591,2592,2594,2597,2599,2602,2607
Seq#:2; N Pos. 2582,2587,2590,2591,2592,2594,2597,2599,2602,2607
Seq#:9; N Pos. 1,2,3,5,9,10,11,12,13,14

Invalid <213> Response:
Use of "Artificial" only as "<213> Organism" response is incomplete, per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:9,10,11